

## EXHIBIT 1

Table 1: SEQ IDs of Claim 29 of Present Application

Mean	33,74
standard deviation (std)	2,5
relative standard deviation ( = std/mean)	
[%]	7,4%
minimum	30
maximum	36,67
range (max-min)	6,67

Table 2: Probes of Thunnissen et al.

SEQ ID NO	Sequence without spacer	GC (%)
24	AGAGGAGCAGGACGACAATG	55
25	TGAAGACGAGGAGGACAATG	50
26	CCATTAAGGTGTCCGAAGC	50
27	AGATGTGTCAAAAGCCAAG	40
28	CGAGGAGGAGCATGGAAACC	60
29	CCATTAACTGTGTCAGAGAC	45
30	ATTGACAGTATCACAAGCTA	35
31	CAGACCTACGTGACCATATA	45
32	ACATGGCATACAGACATTAA	35
33	GAGGAAATGGAAACCTAG	45
34	TAGTAAACGACTTGTGATC	35
35	AGCACTGGAATATCCAGGG	50
36	CTTTATTGTATACAGCCAAA	30
37	AGTAATGGAATTCACATA	30
38	TAGCACATGTTGTCTGATC	40
39	AGAATACTATGAACAAGACA	30
40	AGATGTTCAAGGCTAAAG	35
41	AACATTGGAAACATGTAGAG	35
42	GAAATGTATACGATATGAAT	25
43	ACATGGTATTACCAAACCTAA	30
44	TTTGTTTACAAAGCAAAG	25
45	TTTAGCGCTGAACGACAACG	50
46	TGTTATTACACAAAGCAAAG	30
47	GTTTCTTTACAAGGACGTGG	45
48	AGAGGATCAGGAAGACAATG	45
49	CTATAATGTATACAGCCAGA	35
50	AGACATTAATGAACACATAA	25
51	AGAGGGATCTGATCAACAGG	50
52	CTTGATTTATAAGCTAAA	20
53	AGTTTTTTTCCACCACTTG	35
54	AGAACATTATGAACAGGACA	35
55	AGAGGGACCTGACGAACAGG	60
56	AGTAATGGGAACCCACTATA	40
57	TATATGCACTAAATGATGTA	25
58	TTTGAATTGATCAAGAGG	35
59	AACATTACGAGACTGATAGT	35

Mean	38,61
standard deviation (std)	10,0
relative standard deviation ( = std/mean)	26,0%
[%]	
minimum	20
maximum	60
range (max-min)	40